Atty. Docket No.: PF3623USw
S/N 09/936,506

In the Specification :

On page 5, please replace the paragraph starting on line 7 with the following paragraph :

For the purposes of the present invention ΔG can be calculated using the RNA structural prediction program MFOLD (Zuker M. and Jacobson A.B. (1995) Nucleic Ac. Res. (23) 2791-2798).

- 2 -

Atty. Docket No.: PF3623USw
S/N 09/936,506

On page 26 please replace the paragraph starting on line 20 with the paragraph below

"Sequence identity"

For the purposes of the present invention, sequence identity may be determined, for example, by using the ALIGN program (version 2.0). This calculates a global alignment of two sequences. (See Myers and Miller,(1989) CABIOS, 4, 11-17). Gap penalties: -16/-4.

- 3 -